**Supplemental Figure S3: Eaf7 associates with the coding region of highly transcribed genes.**

(A) Chromatra graphs (Hentrich et al, 2012) showing the enrichment of all labeled probes plotted according to transcriptional frequency (Y axis) and localization throughout the gene (5' to 3' end of gene on the X axis). The transcription start site (TSS) is indicated, as well as the -3, +3 and +10 nucleosomes. The quantile normalized data was binned into segments of 150bp on the X axis. The Y axis is labeled with the 5 classes of transcriptional frequency ((Holstege et al, 1998) Transcriptome updated Feb 2005). (B) Genome-wide average profile obtained from high resolution anti-Myc ChIP-chip as in Figure 5 but from an untagged strain. (C-D) Examples of anti-Myc signals from the untagged strain at the six specific loci showing Epl1 and Eaf7 enrichment in figure 5E-F. Average signals are based on at least two independent experiments.